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↓

Rat	MKLVNIWLLLLLVLLCGKKHLGDRLGKKAFAKAPCPSCSHLTLKVEFSSTVVEYIYAFNGYFTAKARNSFISS	75
Mouse	ST V R TR L	
Human	E S G	

Rat	ALKSSEVDNWRIIPRNNPSSDYPSDFEVIQIKEKQKAGLLTLEDHPNIKRVTPQRKVFRSLKFAESDPVPCNET	150
Mouse	E N	
Human	Y T	

Rat	RWSQKWQSSRPLKRASLSLGS GFWHATGRHSSRLLRAIPRQVAQTLOADVLMQMGYTGANVRVAVFD TGLSEKH	225
Mouse		
Human	R	

Rat	PHFKNVKERTNWTNERTLDDGLGHGT FVAGVIASMRECQGFAPDAELHIFRVFTNNQVSYSWFLDAFNAILKK	300
Mouse		
Human		

Rat	MDVLNLSIGGPDFMDHPFVDKVVWELTANNVIMVSAIGNDGPLYGTLLNPADQMDVIGVGIDFEDNIARFSSRGM	375
Mouse		
Human	I	

Rat	TTWELPGGYGRVKPDIVTYGAGVRGSGVKGCCRALSGT SVASPVVAGAVTLLVSTVQKRELVPASVKQALIASA	450
Mouse		
Human	M M	

Rat	RRLPGVNMFEQHGKLDLLRAYQILSSYKPQASLSPSYIDLTECPYMWPYCSQPIYYGGMPTIVNVTILNGMGVT	525
Mouse		
Human	N V	

FIG. 1A

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Rat	GRIVDKPEWRPYLPQNGDNIEVAFSYSSVLWPWSGYLAISISVTKKAASWEGIAQGHIMITVASPAETELKNGAE	600
Mouse		HS
Human	D Q V S	

Rat	HTSTVKLPIKVKIIPPPRSKRVLWDQYHNLRYPPGYFPRDNLRMKNPLDWNGDHVHTNFRDMYQHLRSMGYFV	675
Mouse		
Human	Q I	

Rat	EVLGAPFTCFDATQYGTLLMVDSEEEYFPEEIAKLRRDVDNGLSLVVFSDWYNTSVMRKVKFYDENTRQWWMPDT	750
Mouse	L I	
Human	S I	

Rat	GGANVPALNELLSVWNMGFSDGLYEGEFALANHDMYASGCSIARFPEDGVVITQTFKDQGLEVLKQETAVVDNV	825
Mouse	I V K E	
Human	I V K E	

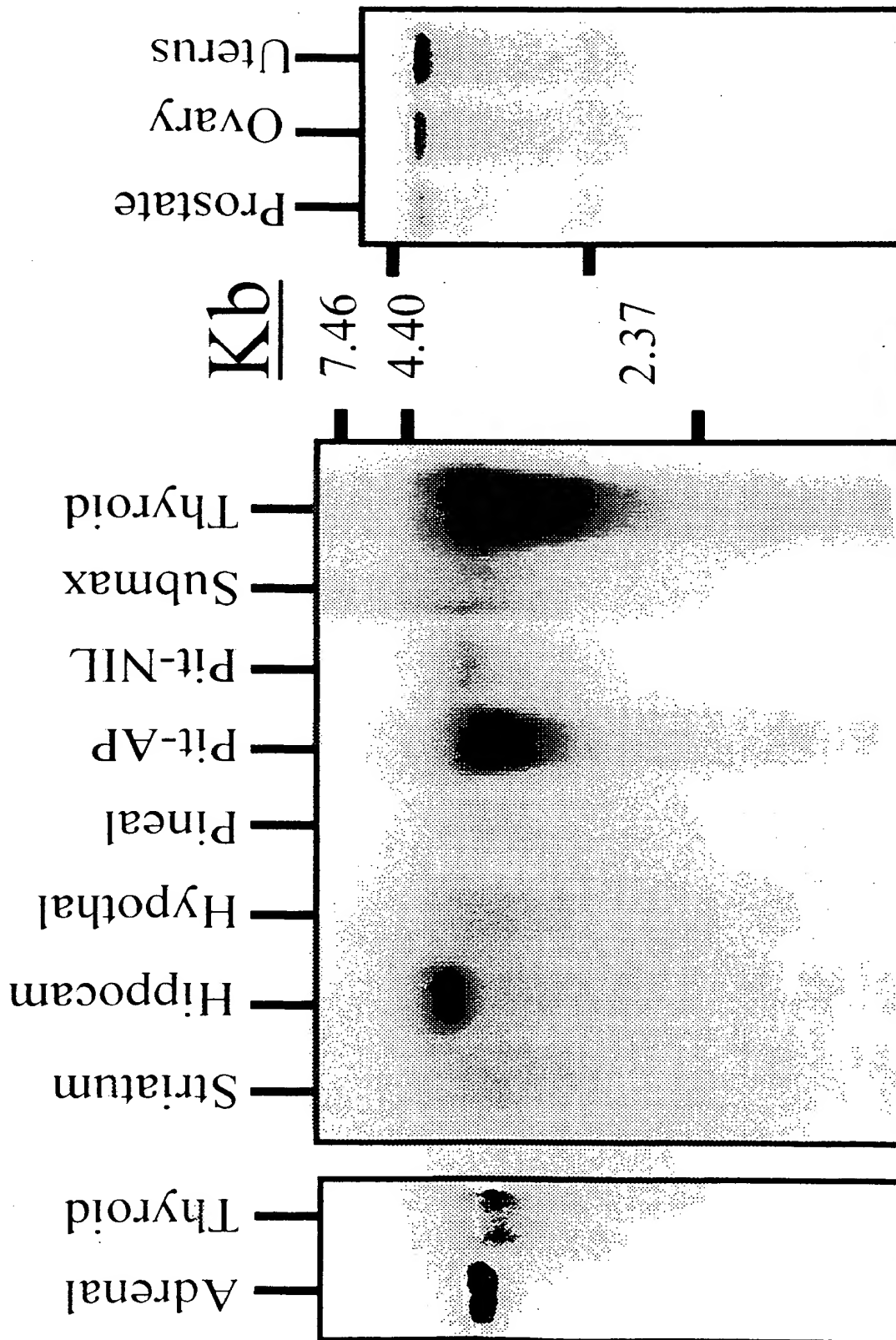
Rat	PILGLYQIPAEGGGRIVLYGDSNCLDDSHRQKDCFWLLDALLQYTSYGVTPPSLSHSGNRQPPSGAGLAPPERM	900
Mouse	S	
Human		SVT

Rat	EGNHLHRYSKVLEAHLGDPKPRPLPACPHLSWAKPQPLNETAPSNLWKHQKLLSIDLDKVVLNFRSNRPQVRPL	975
Mouse		
Human	R	

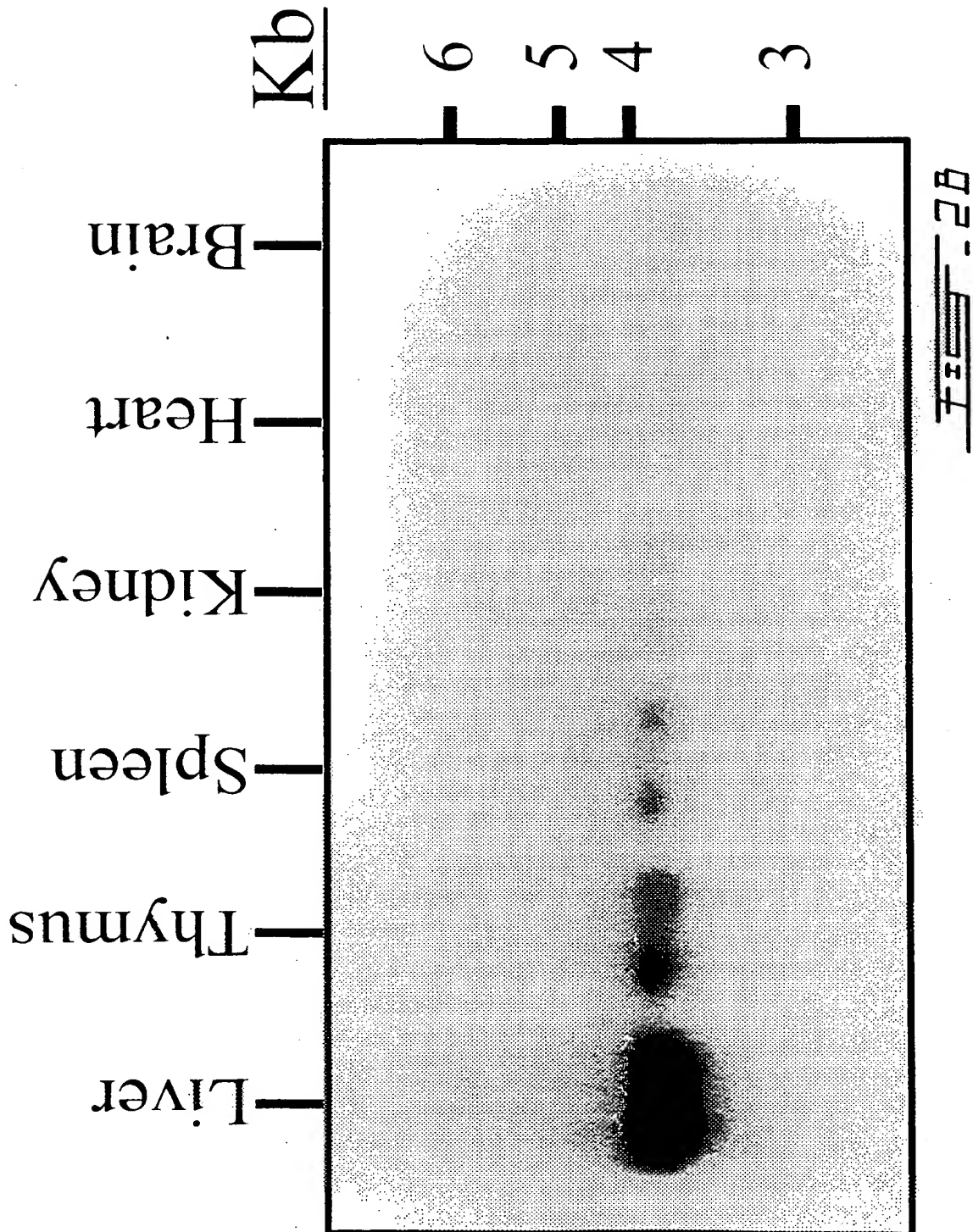
Rat	SPGESGAWDIPGGIMPGRYNQEVGQTIPVFAFLGAMVALAFFVVQISKAKSRPKRRRPRAKRPQLAQQAHPARTPSV	1052
Mouse		
Human	V N K V M V PK	

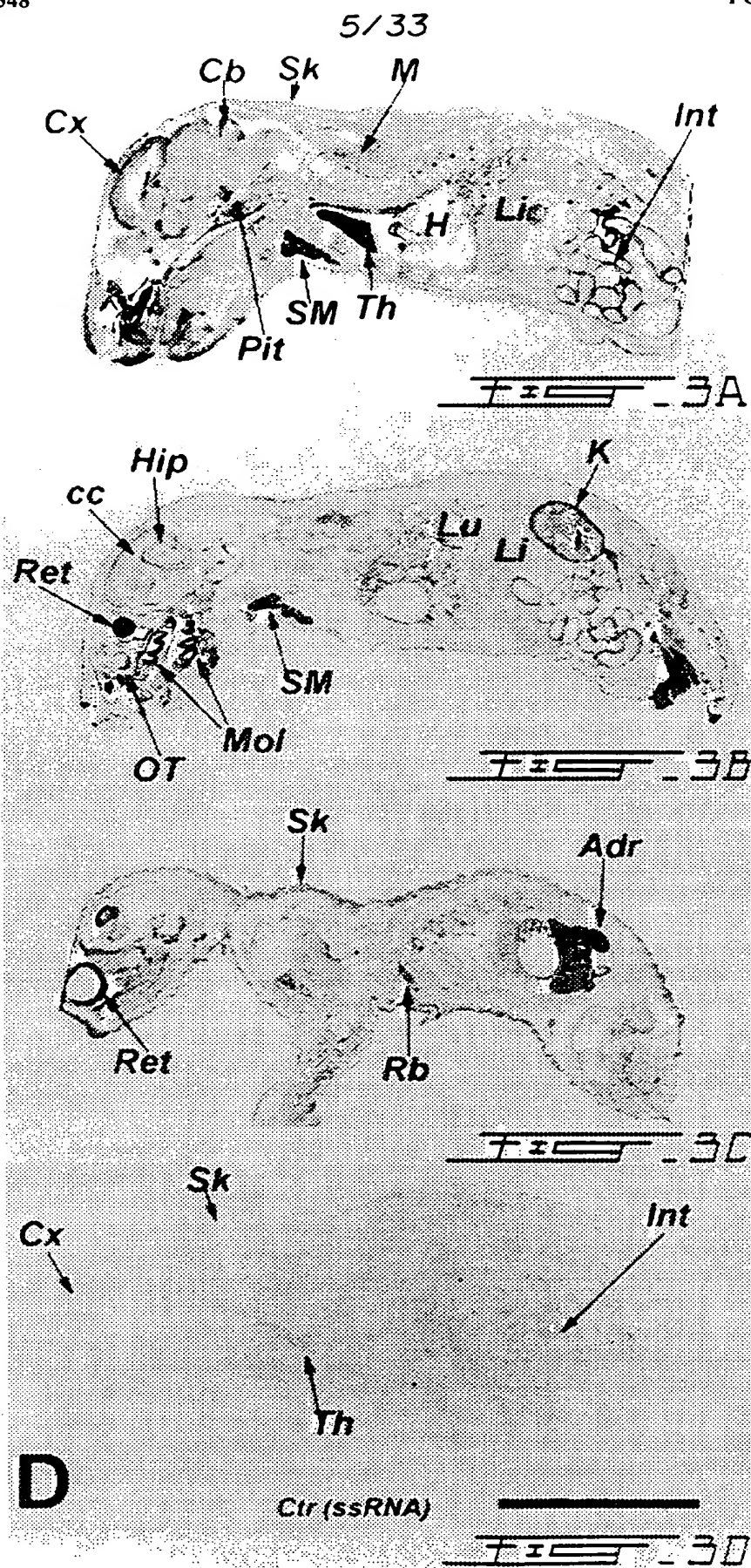
715 - 1B

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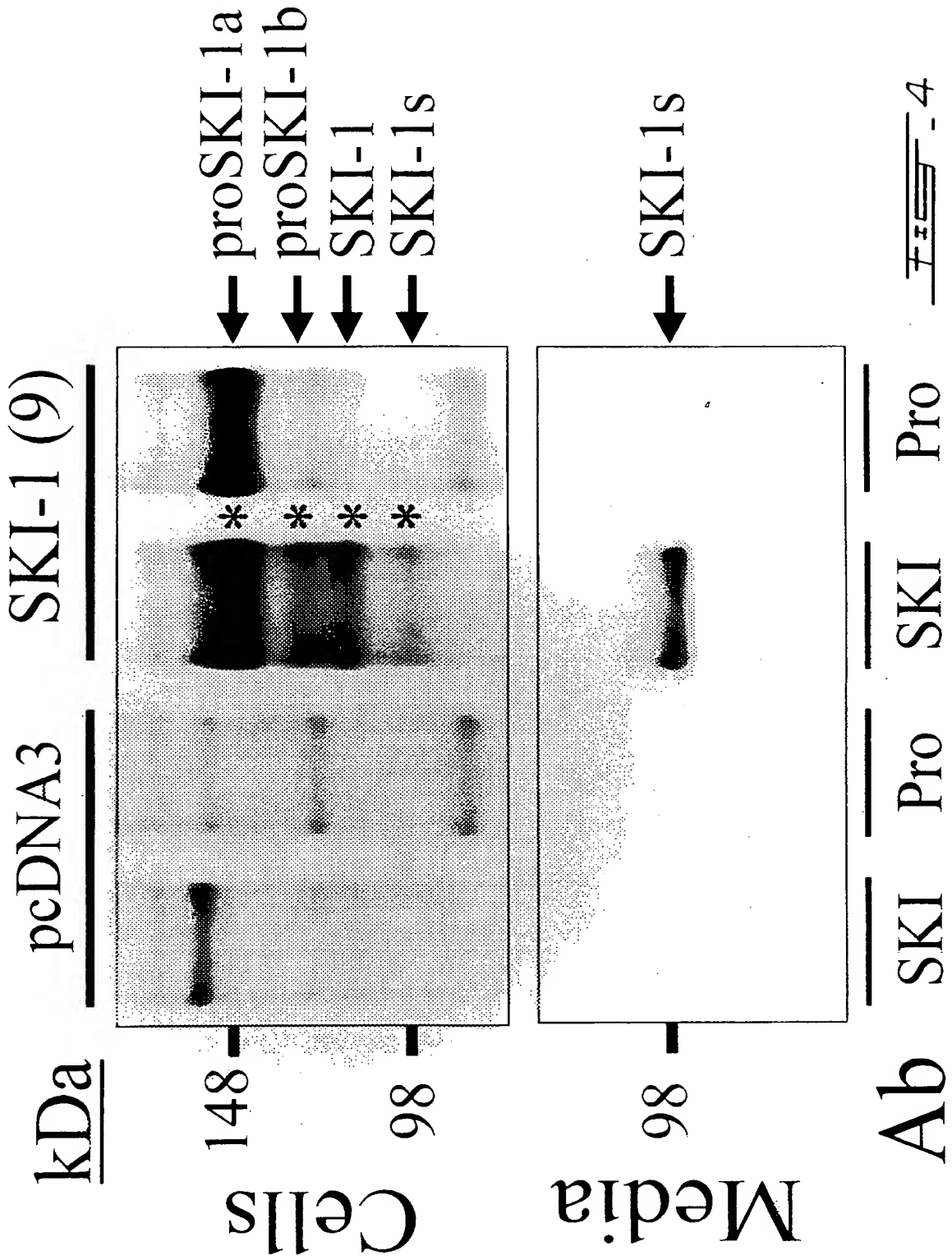


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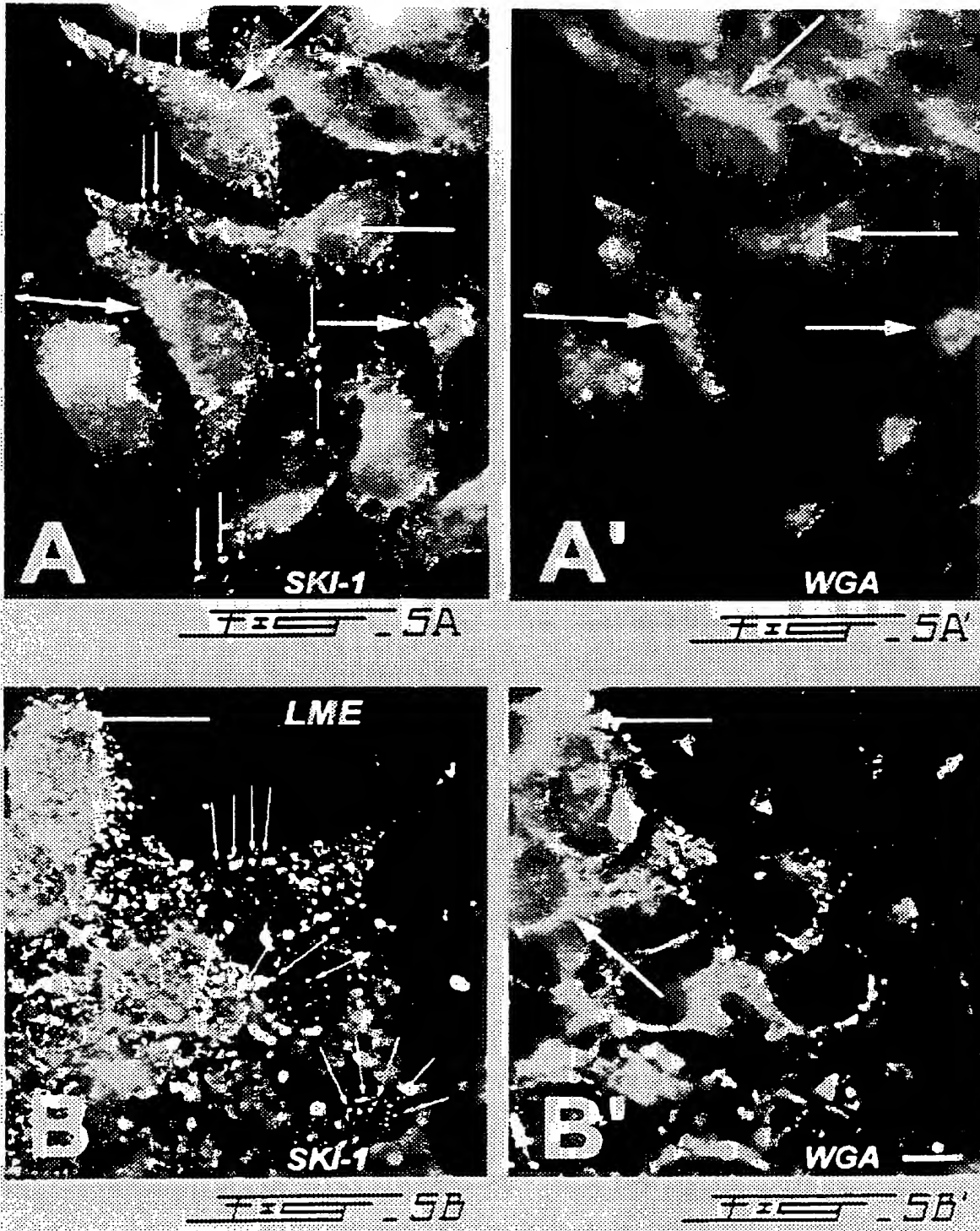




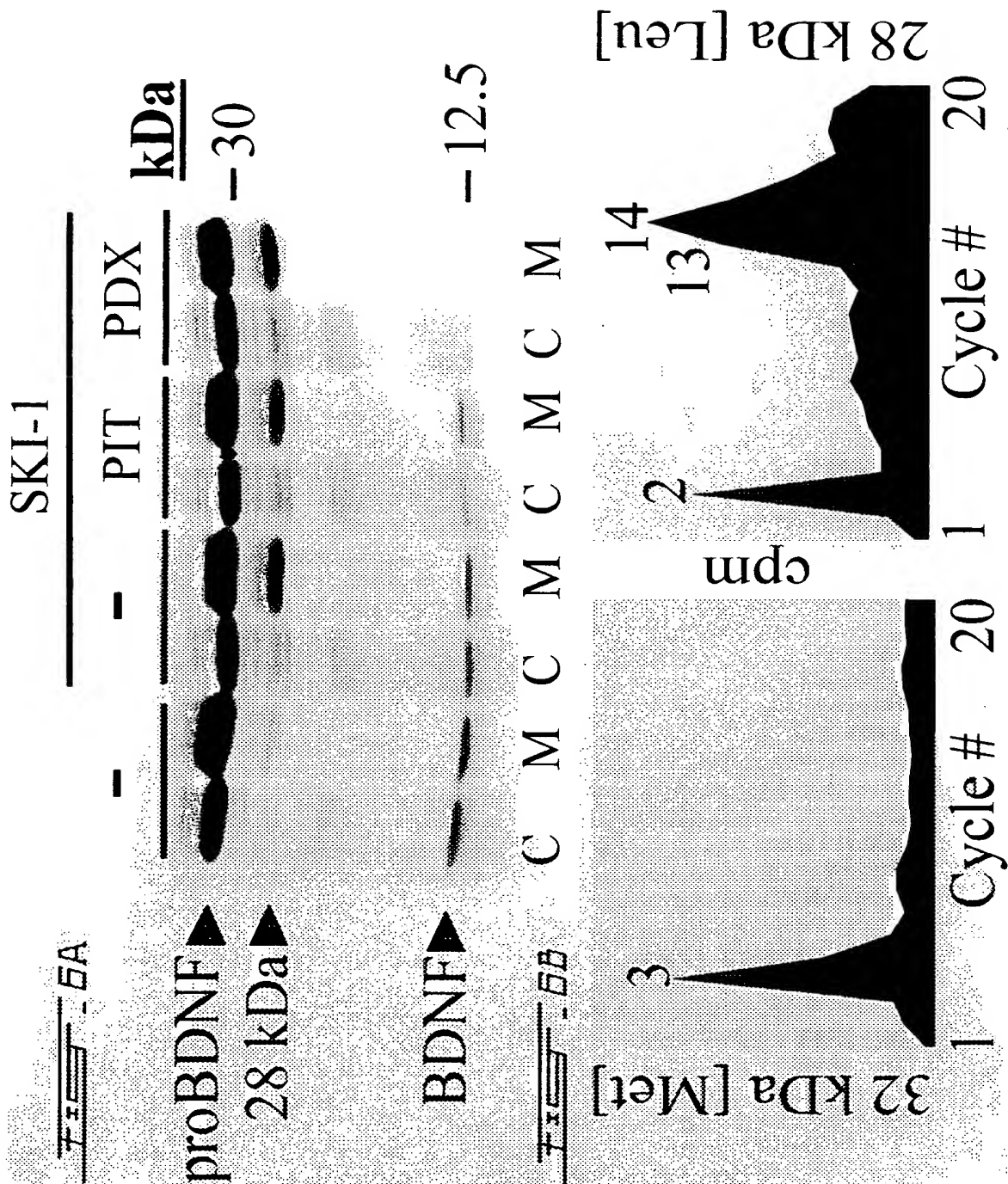
6/33



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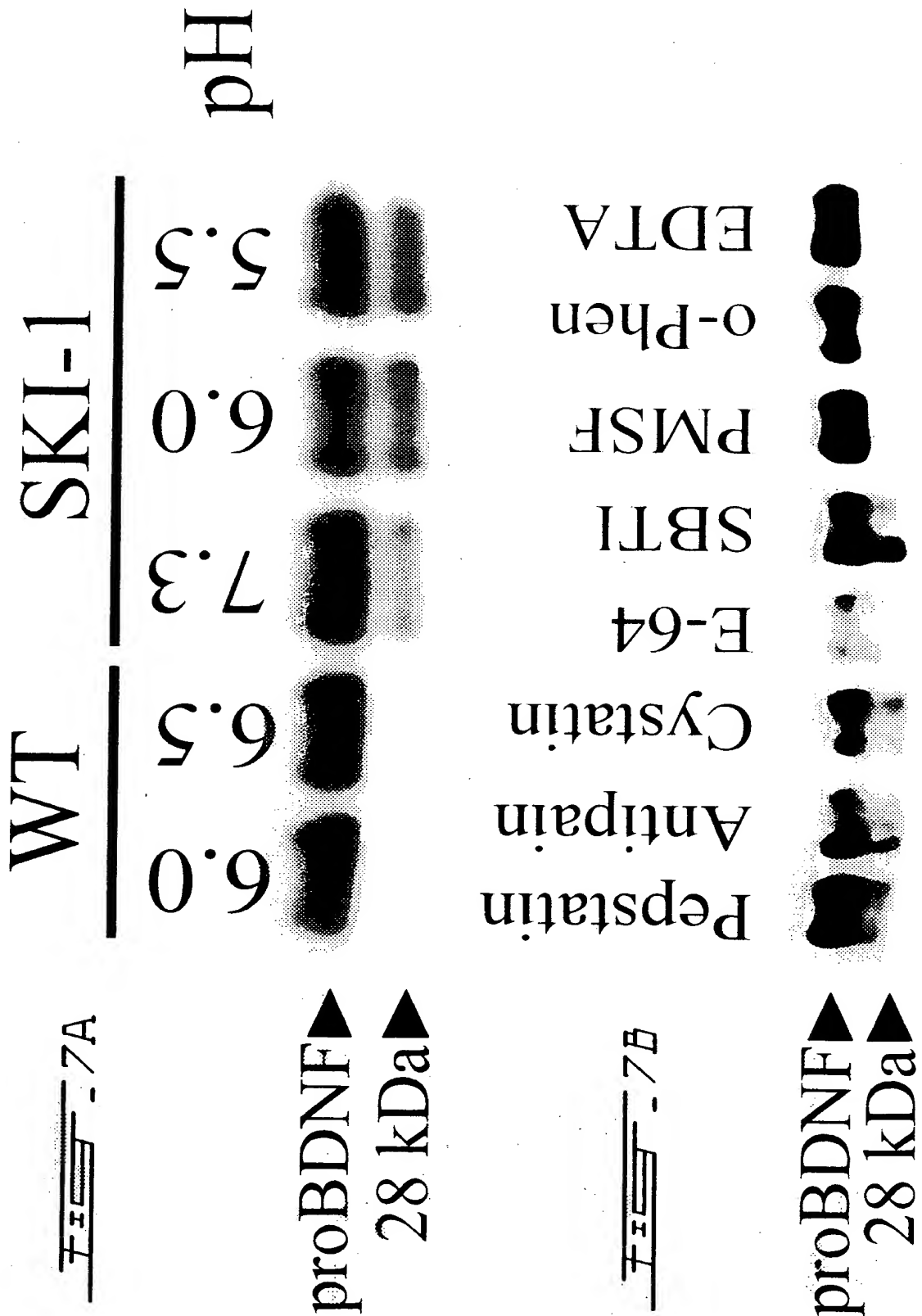


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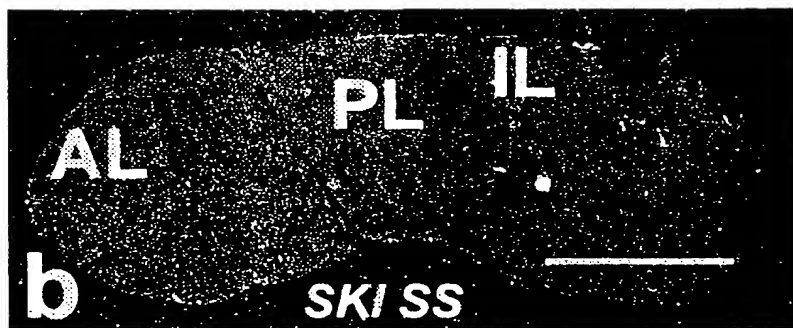




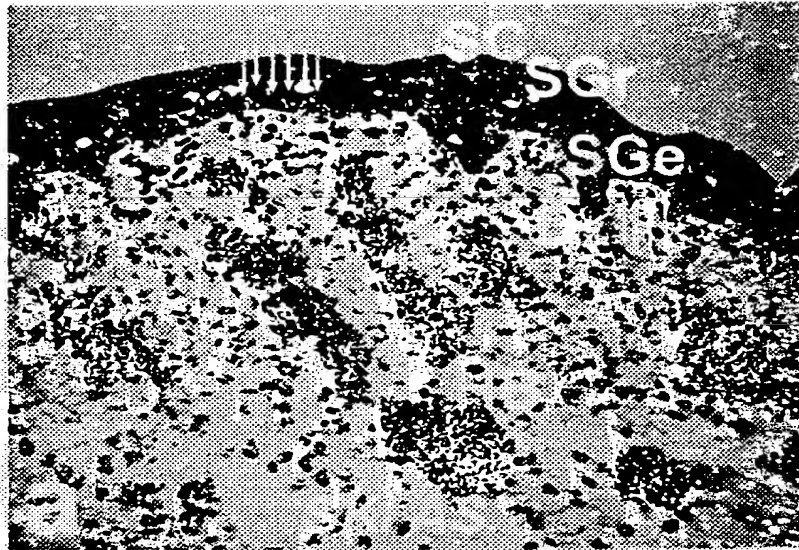
9/33



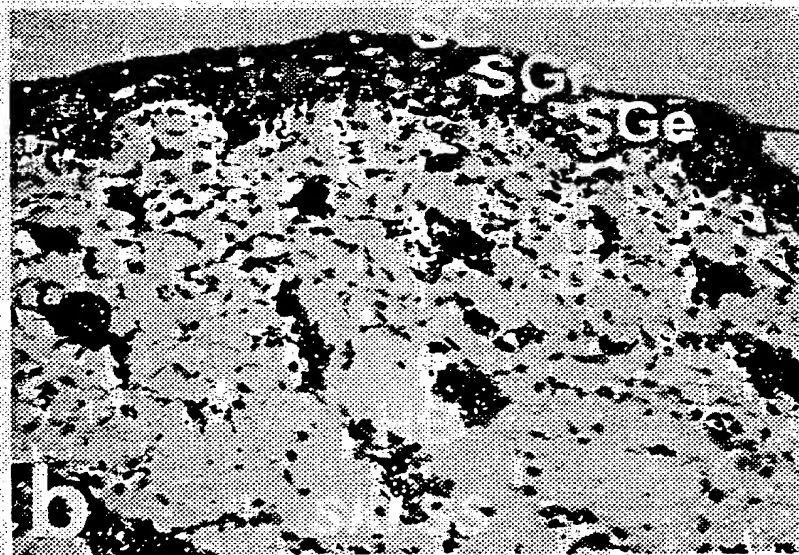
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11/33



T 15 9a



T 15 9b

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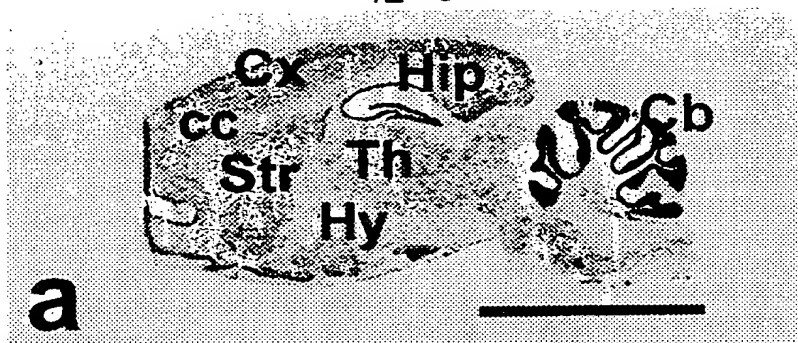


FIG. 10a

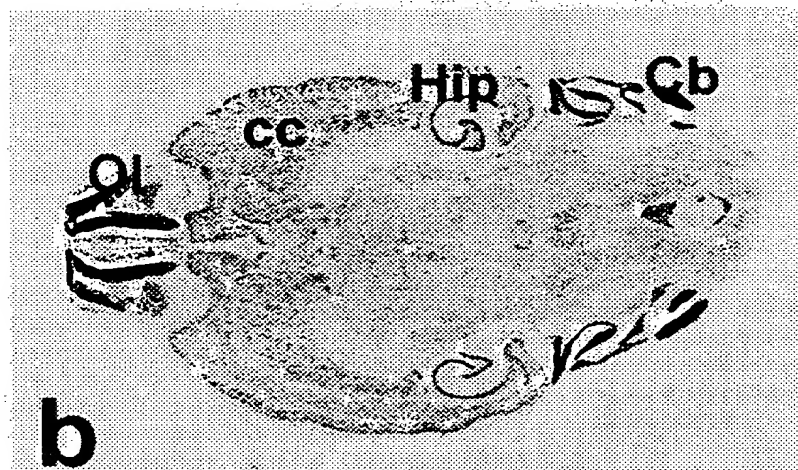


FIG. 10b

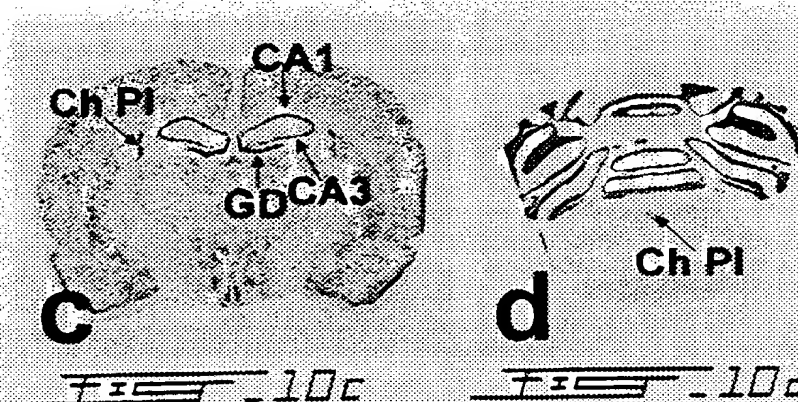


FIG. 10c

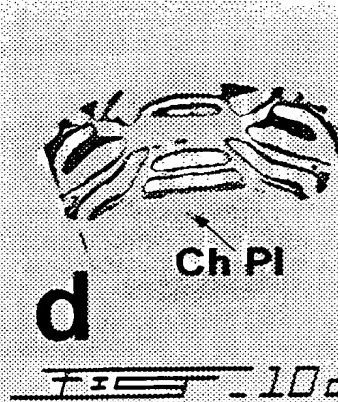


FIG. 10d

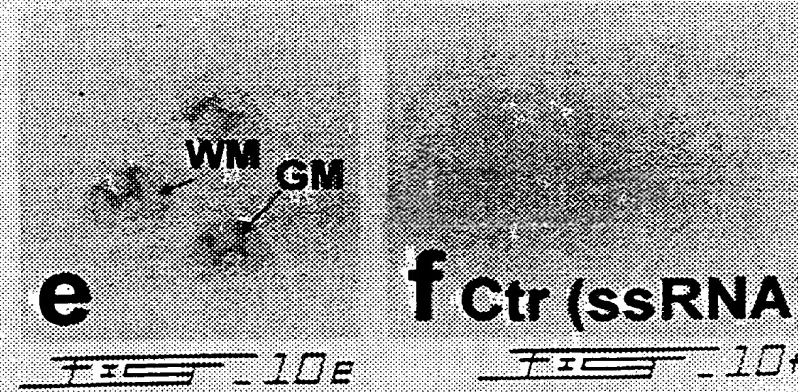
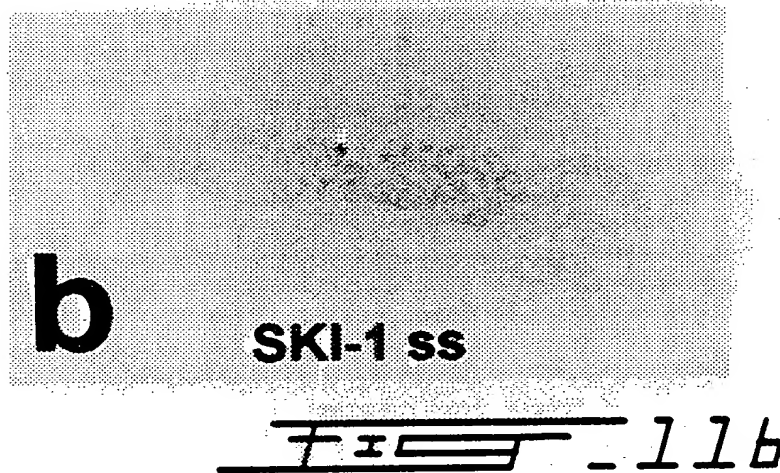
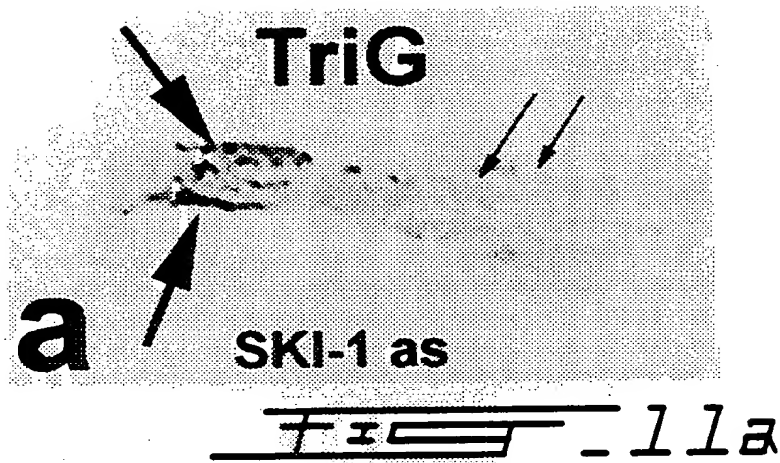


FIG. 10e

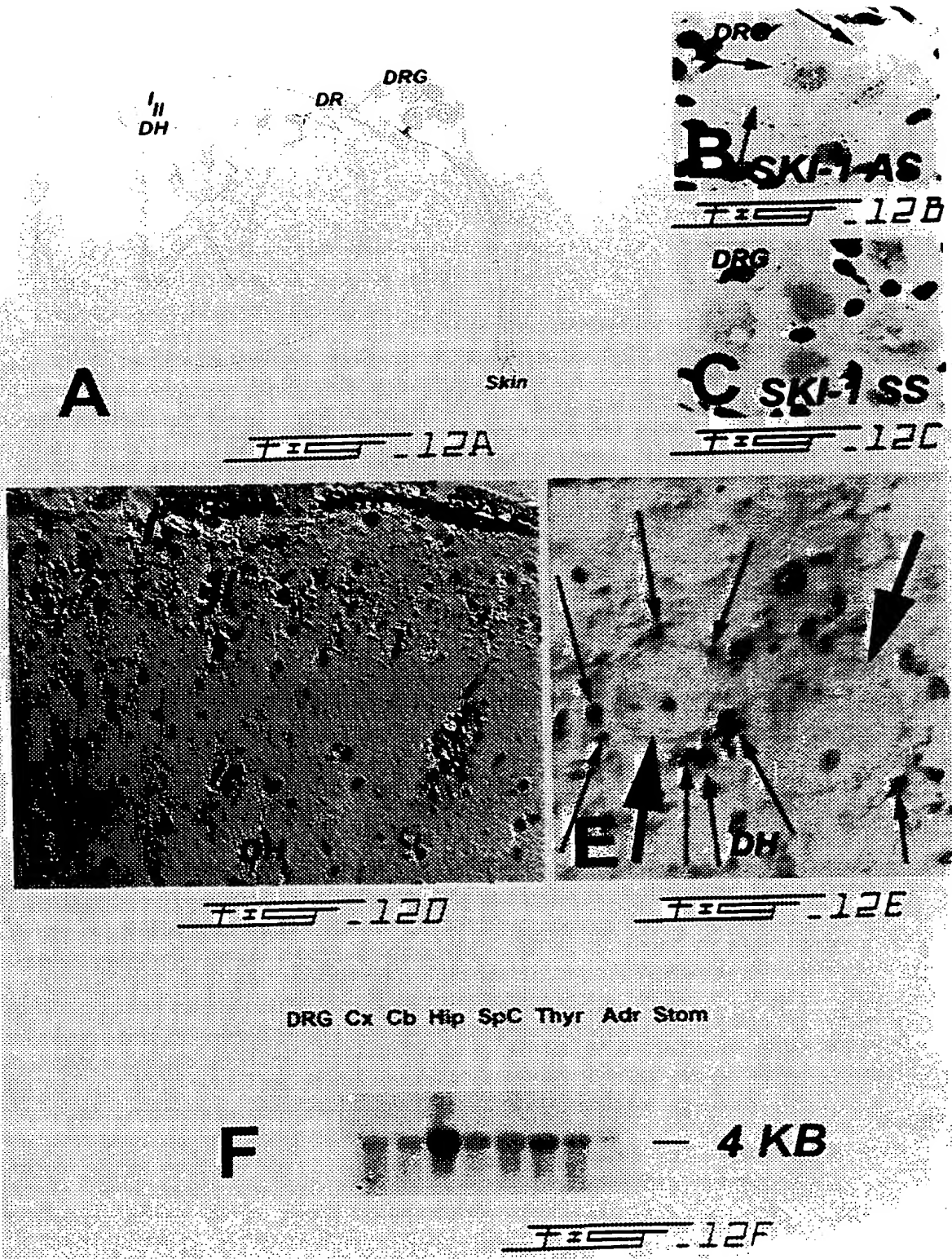


FIG. 10f

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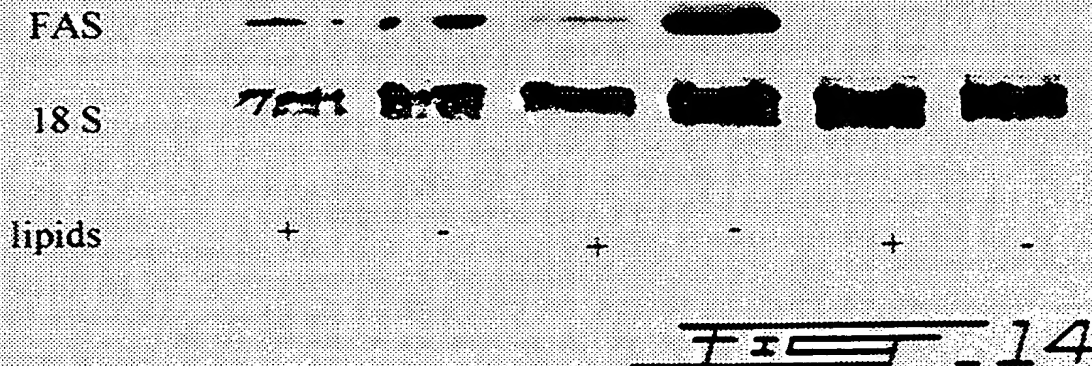
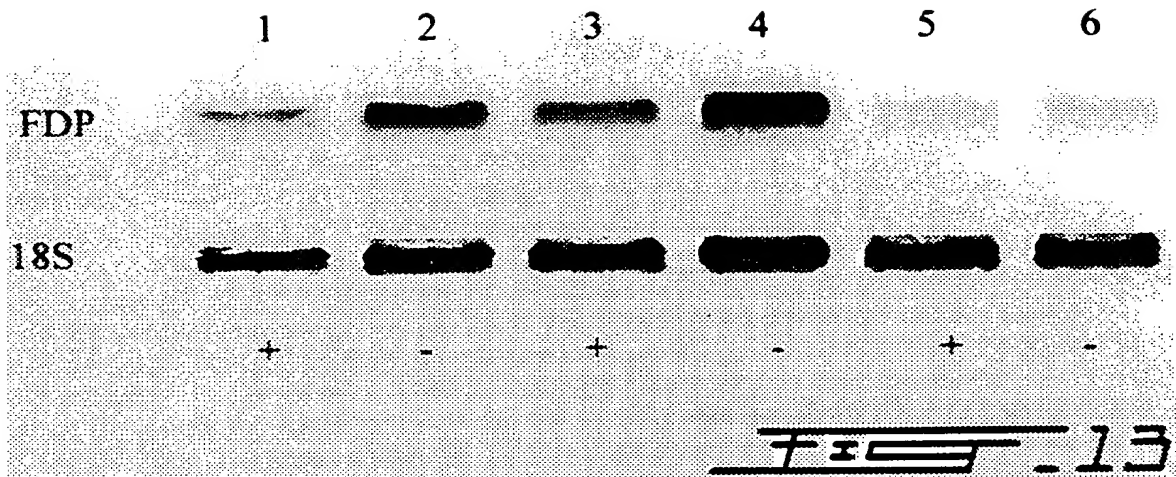


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1	2	3	4	5	1	2	3	4	5

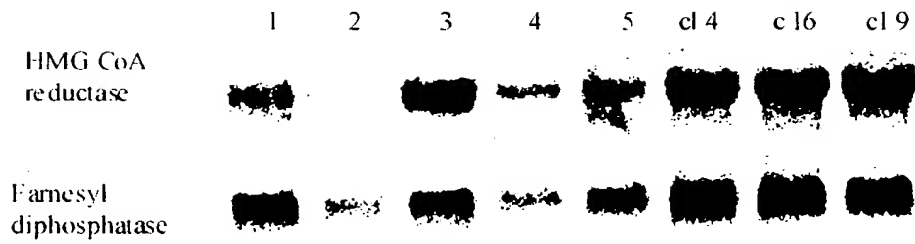
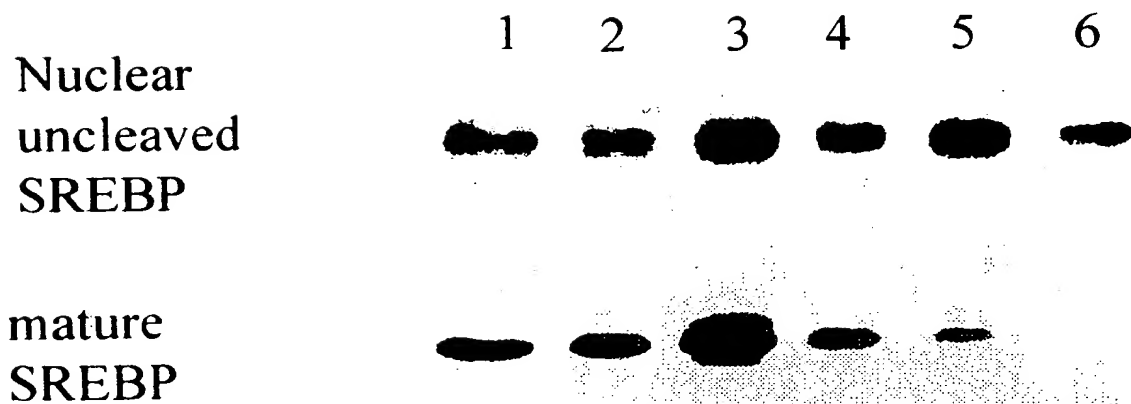
HMG CoA  
reductase

18 S

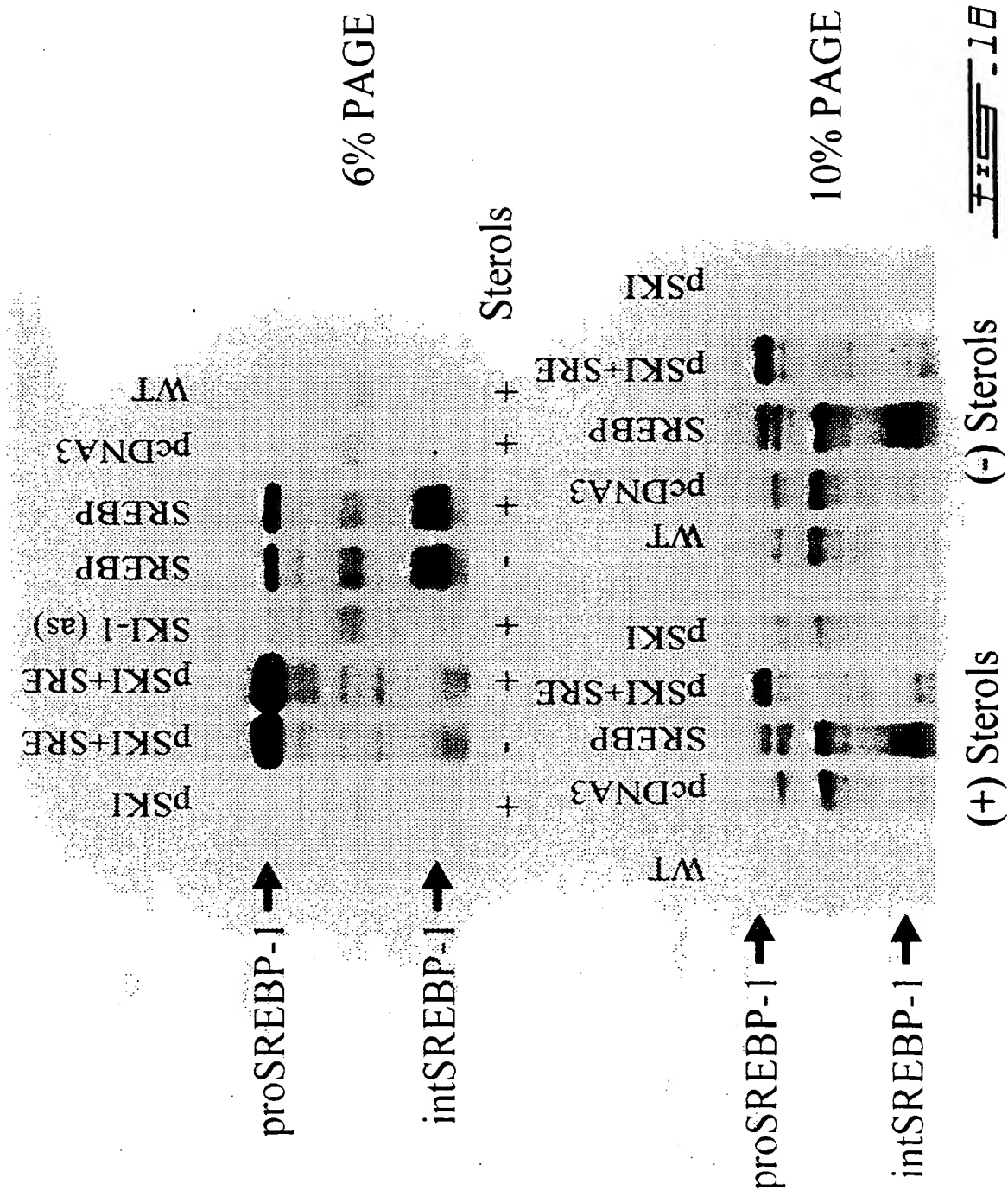
Fig. 15A      Fig. 15B



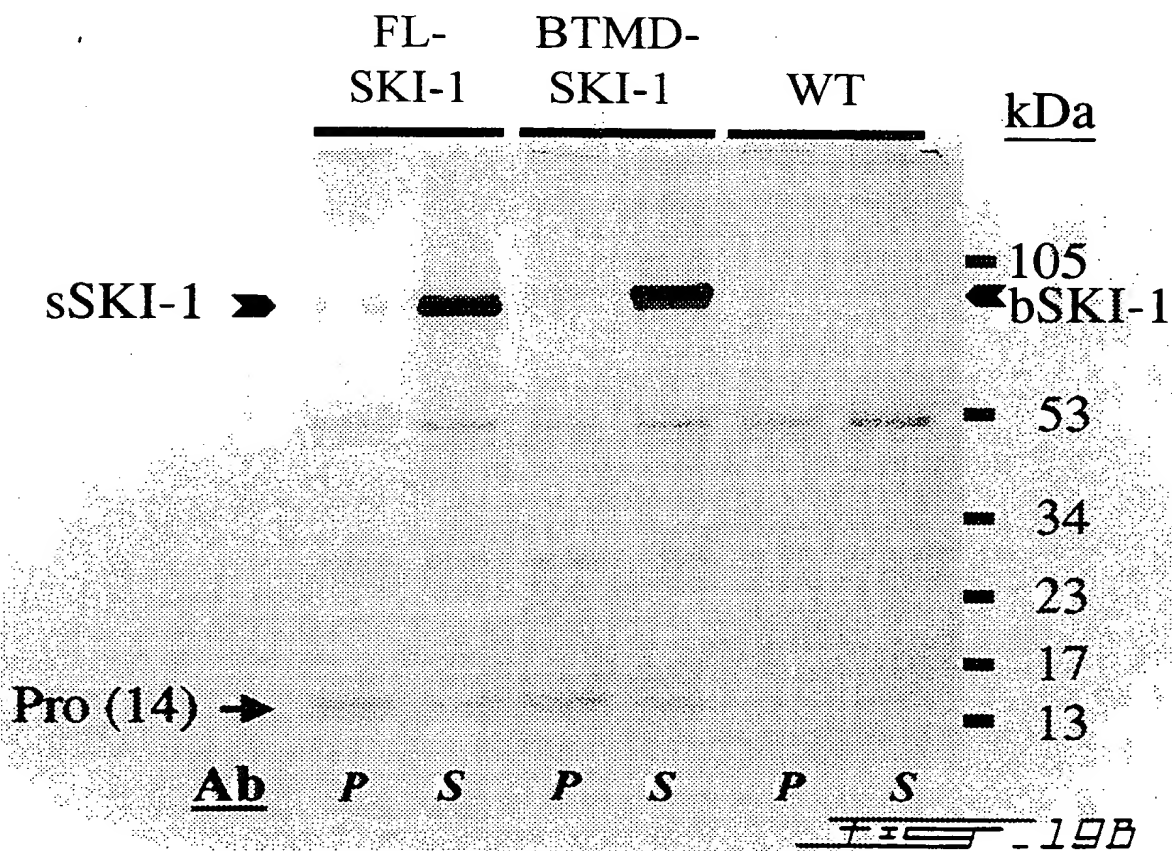
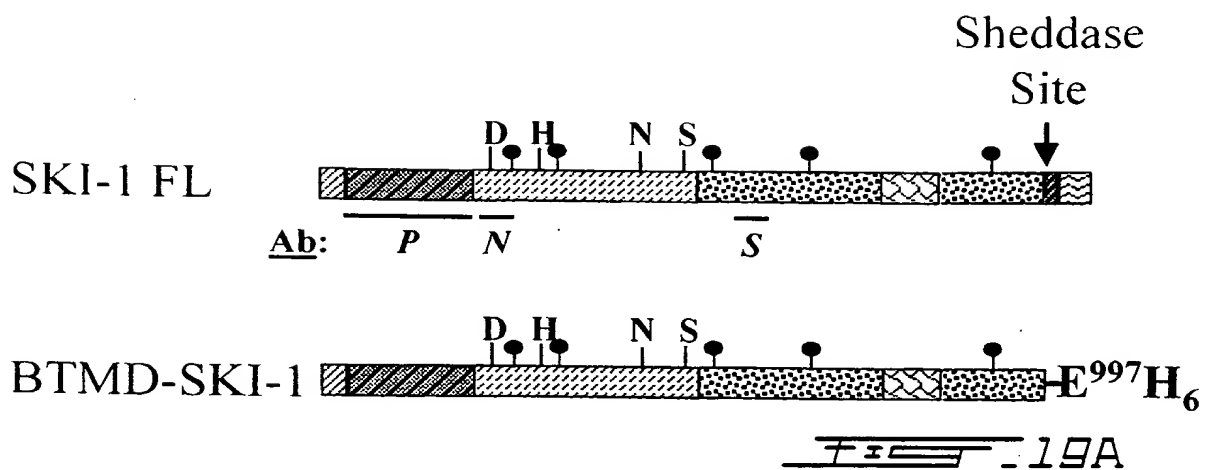
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FIG. 16FIG. 17

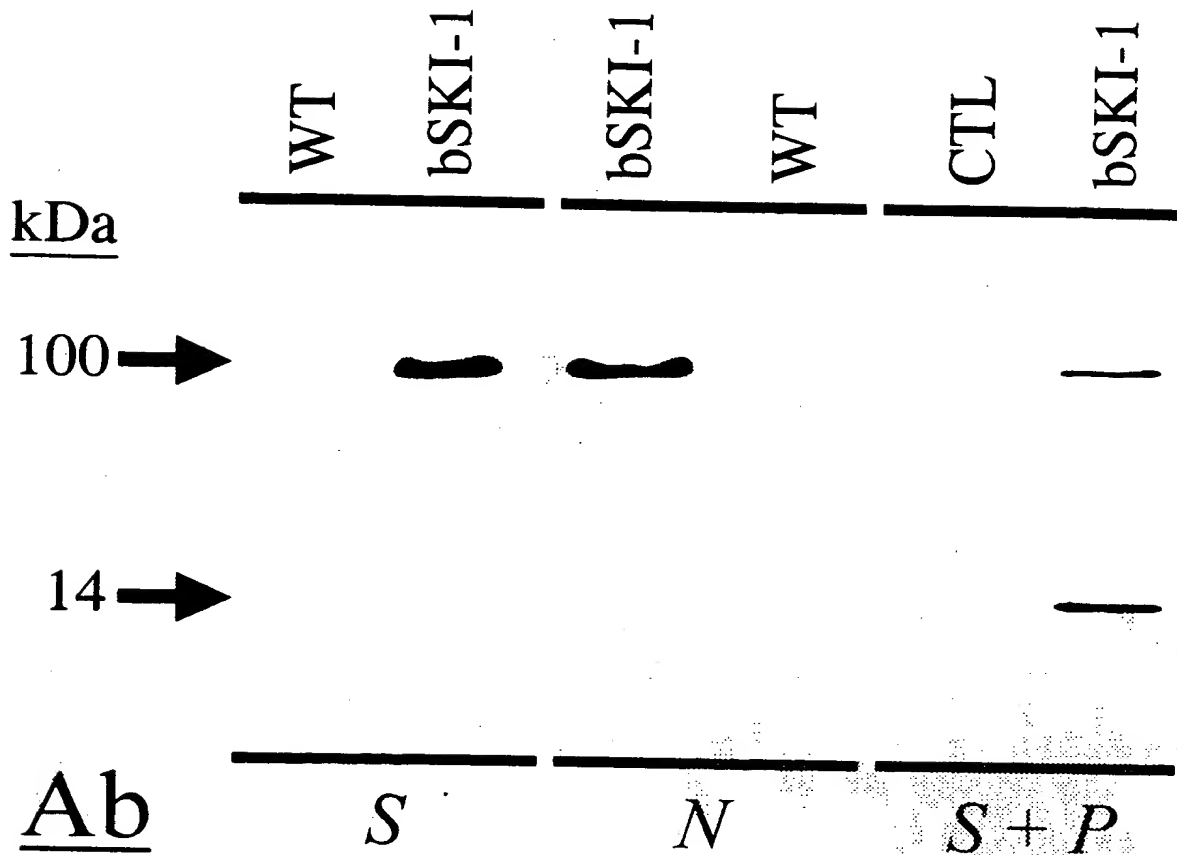
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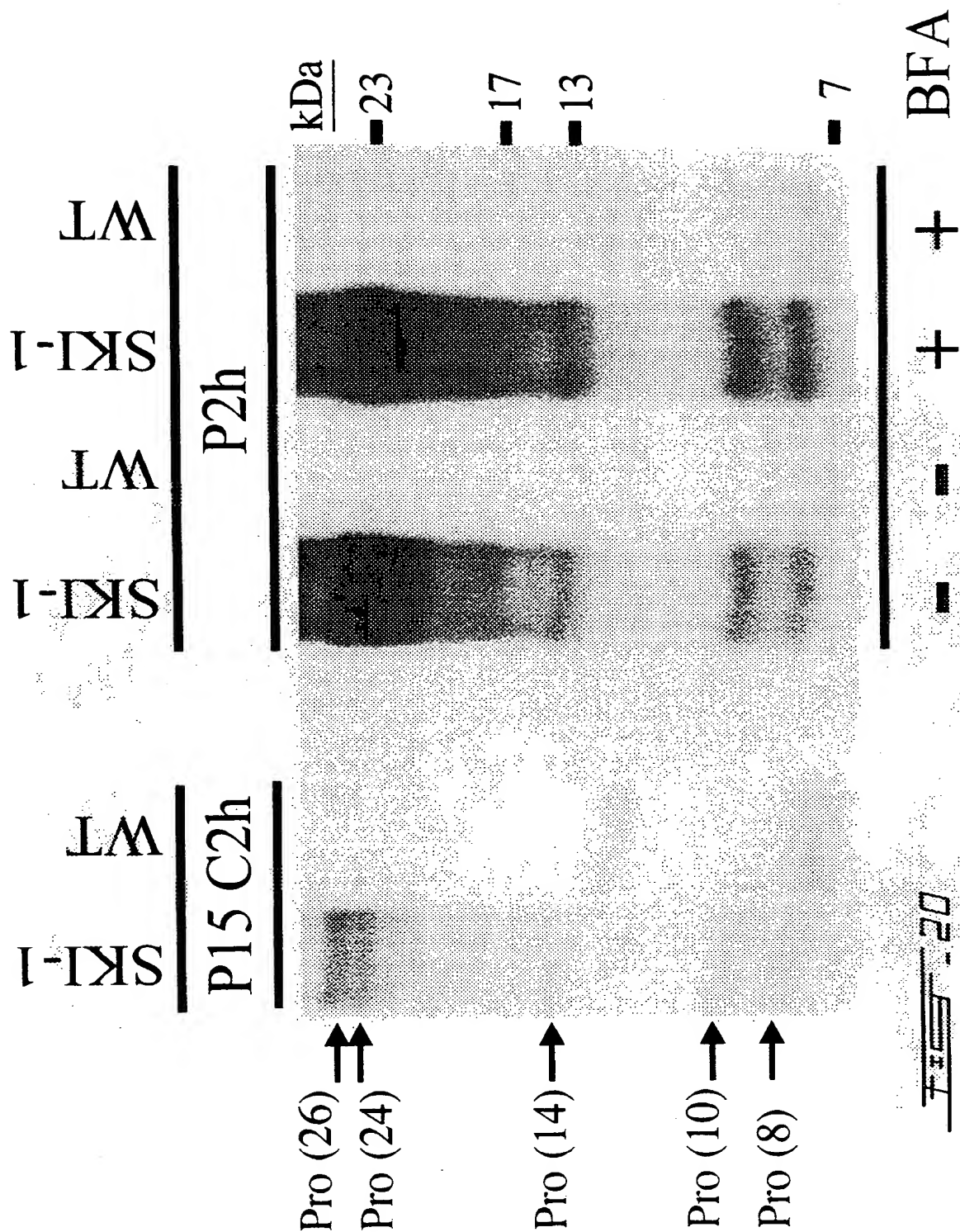


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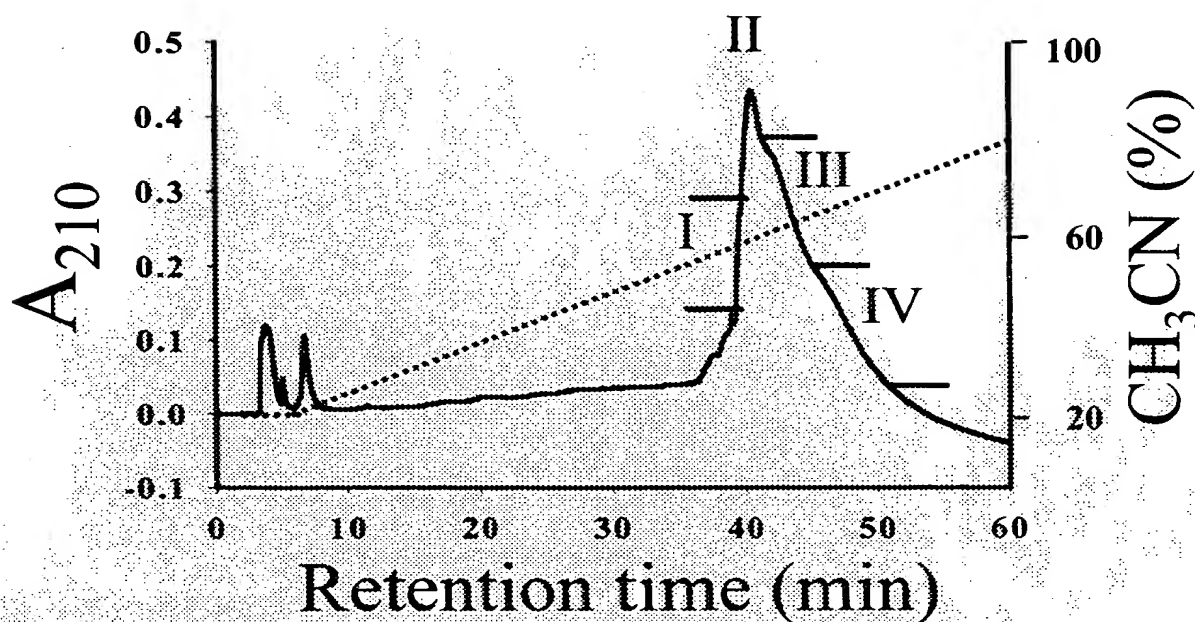


F15-19C

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21A

kDa

I

II

III

IV

24 →

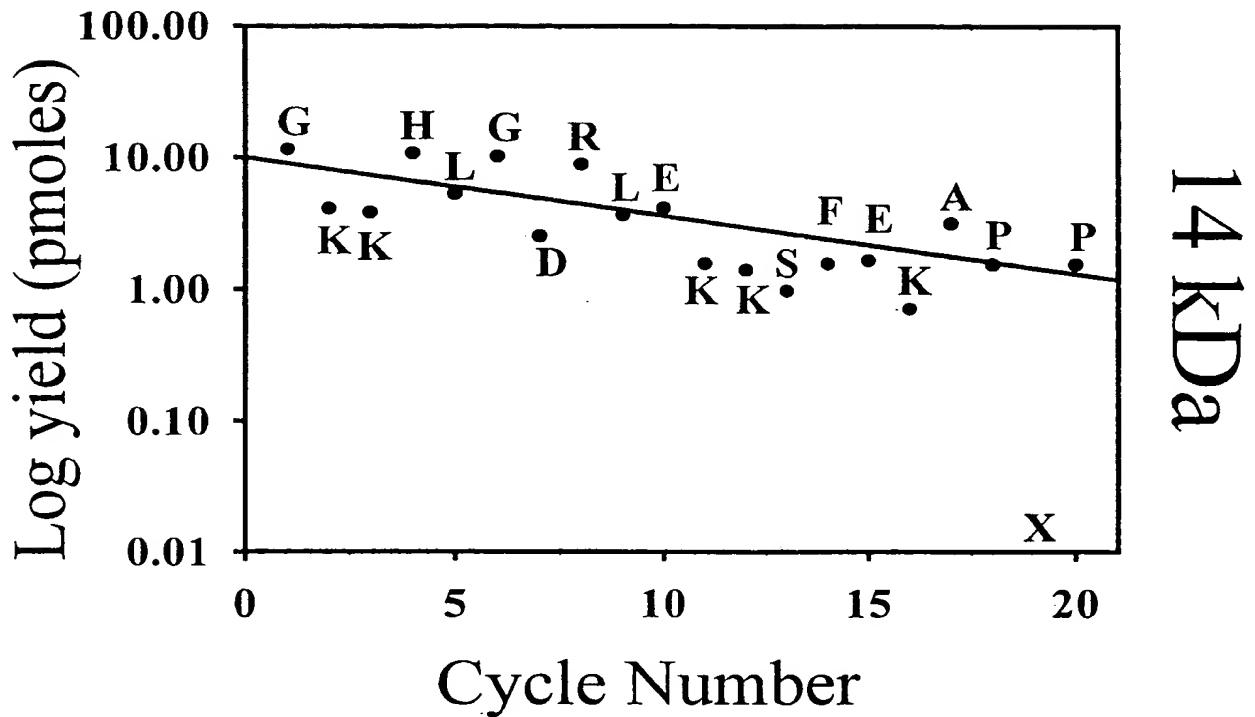
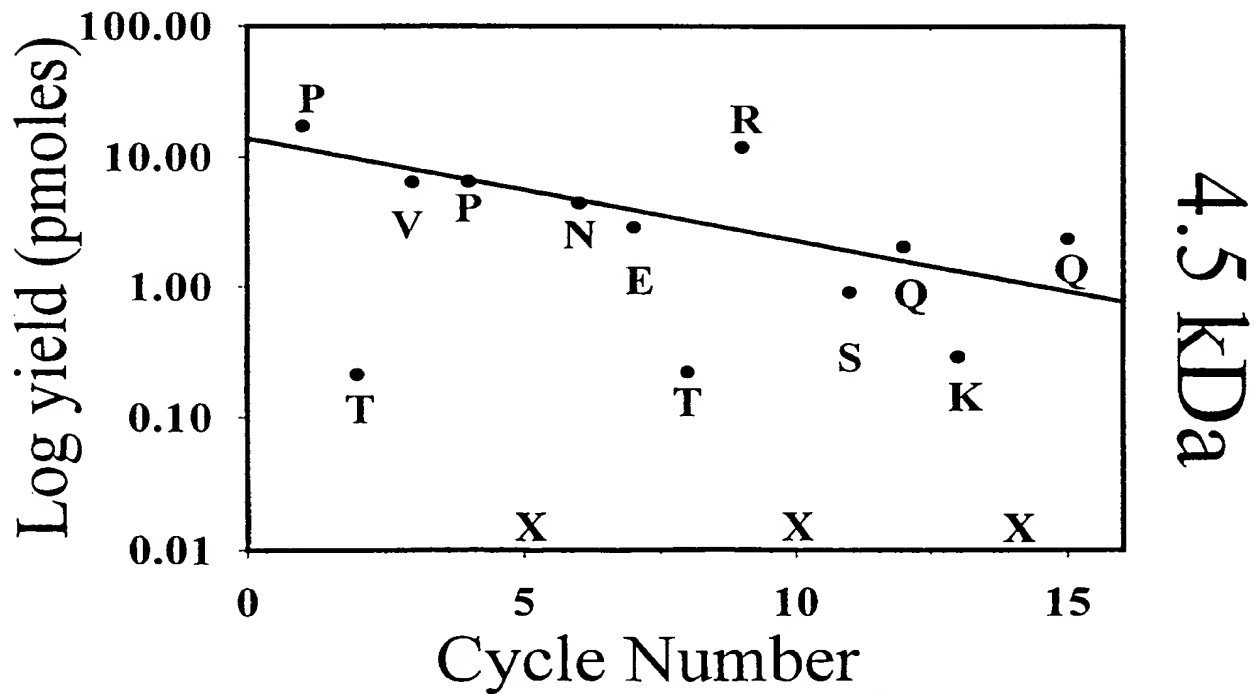
14 →

10 →

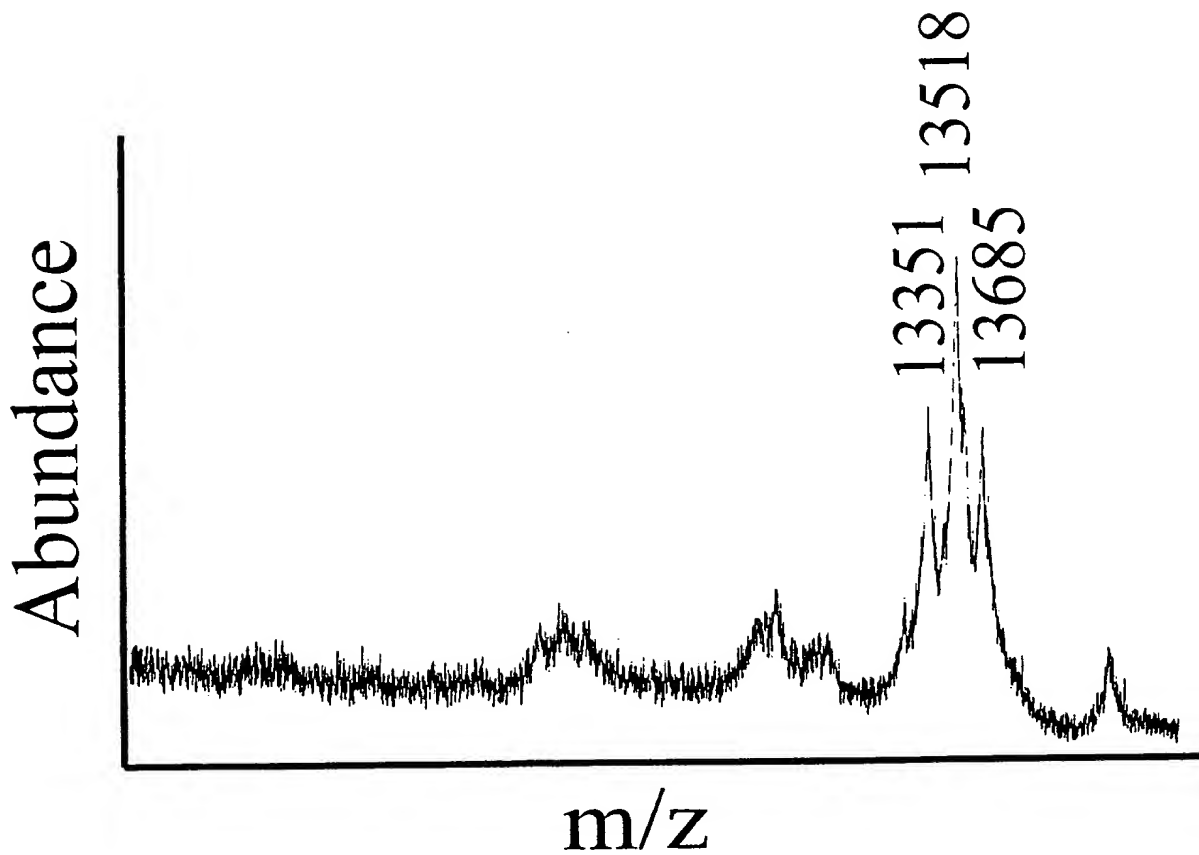
4.5 →

21B

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FIG. 21CFIG. 21D

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QRKVF↓R↓SL↓KYAE

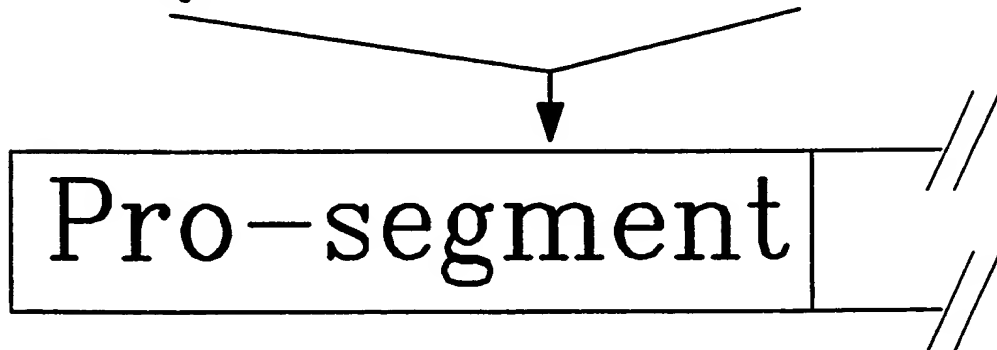
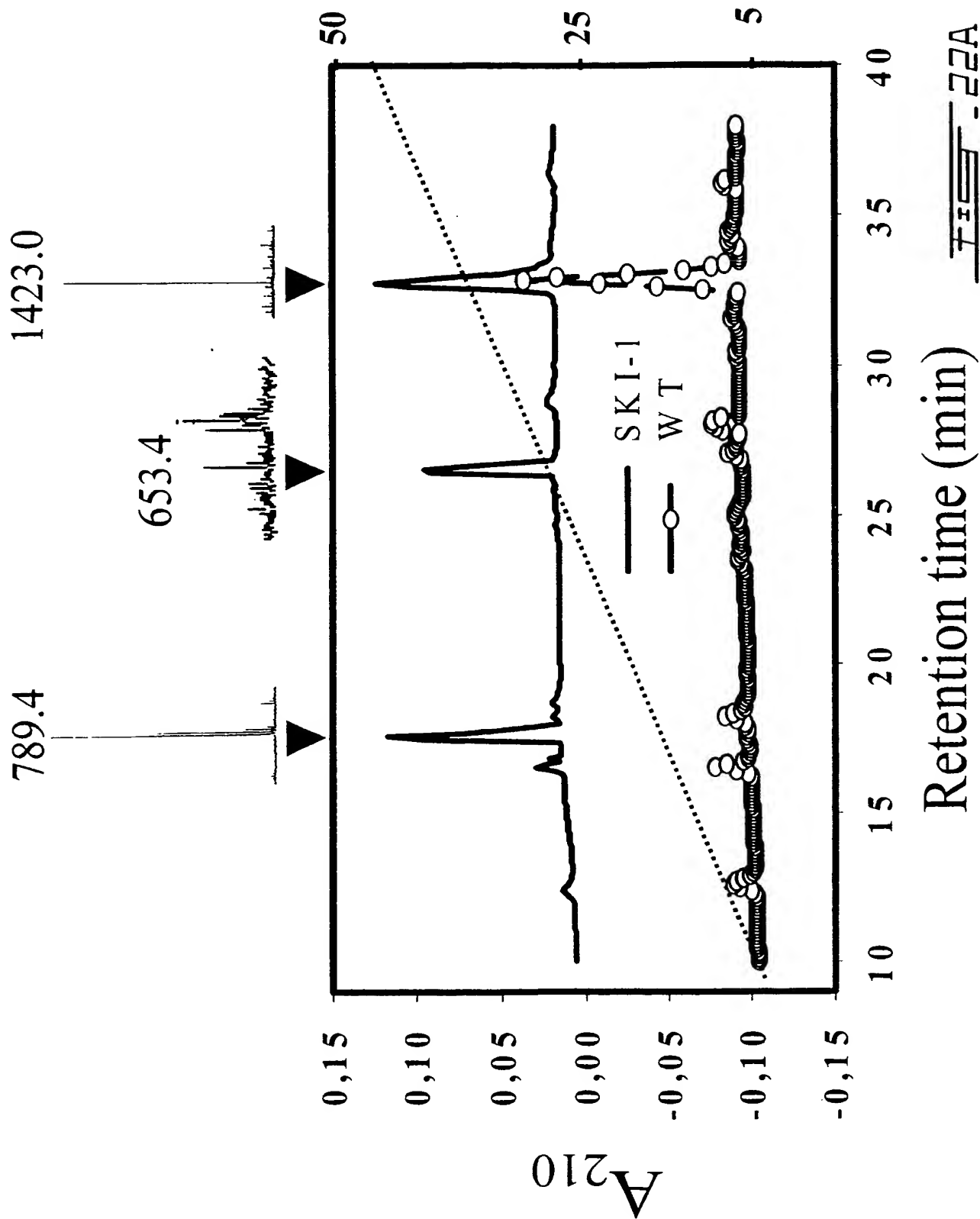
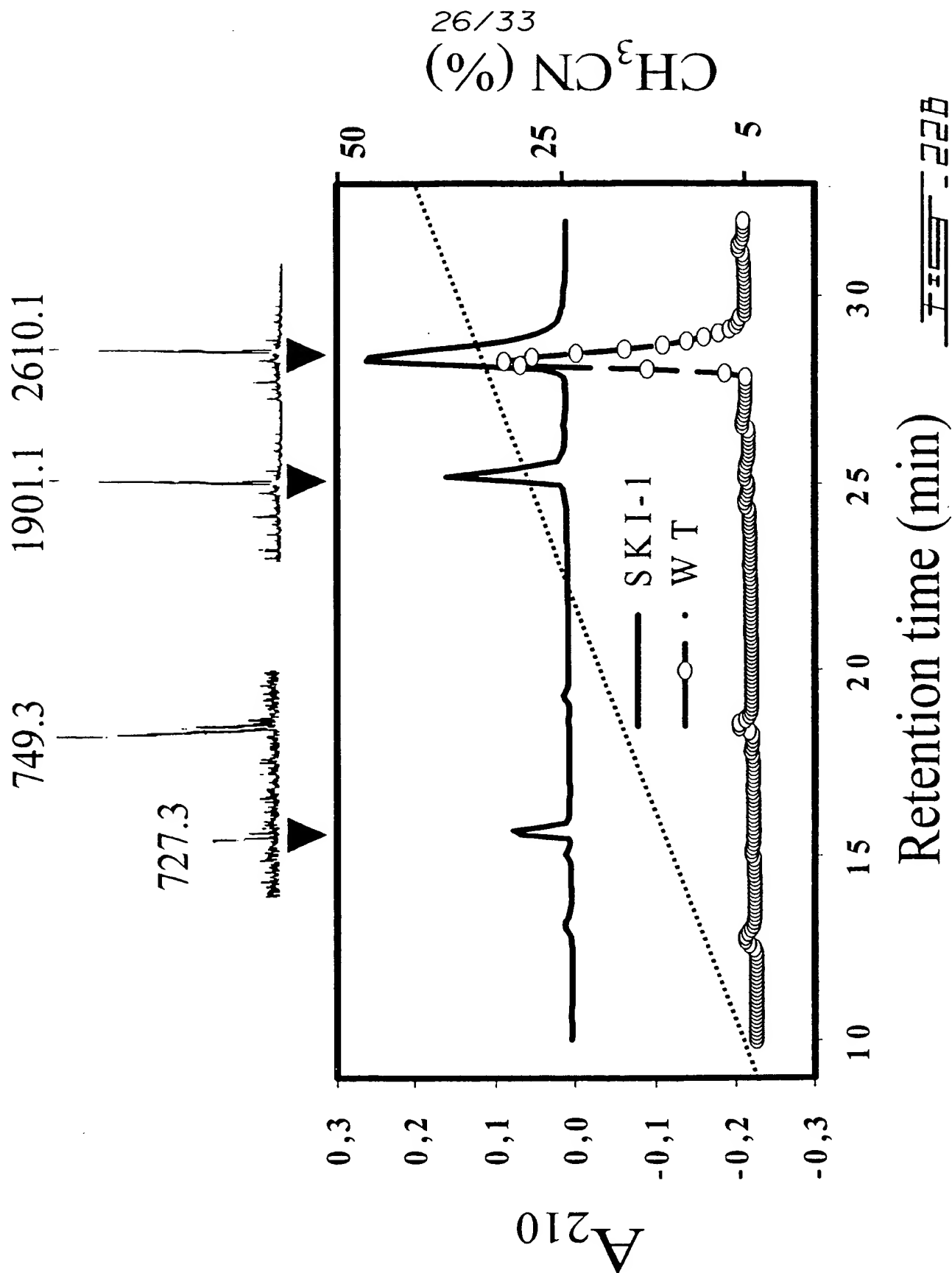


FIG. 21E



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(%)  $\text{CH}_3\text{CN}$





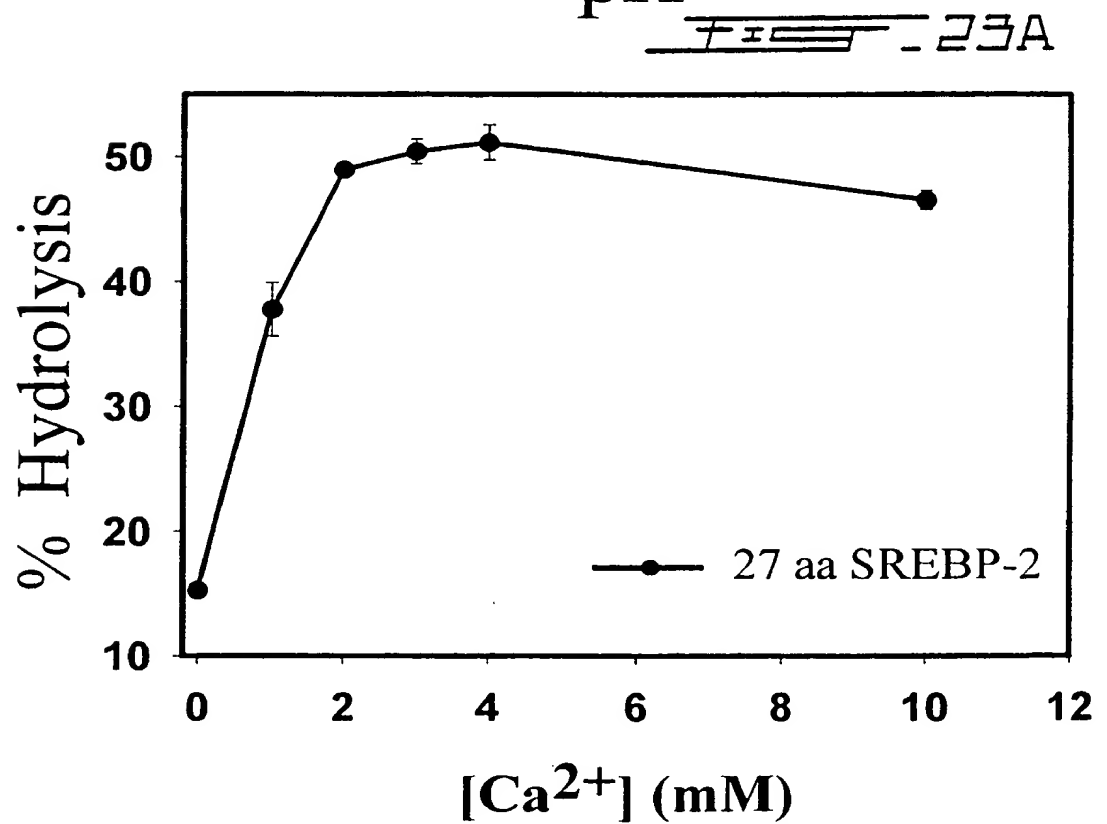
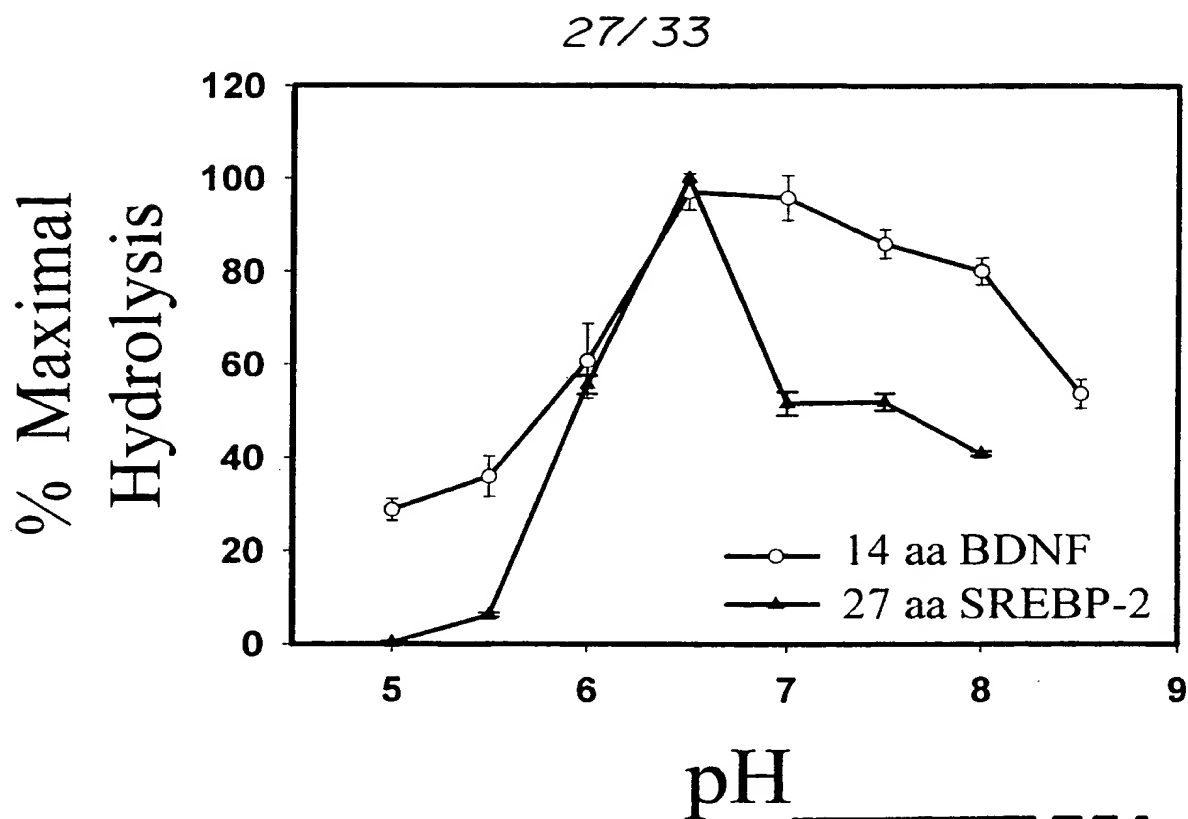


FIG. 23B

DPKK-[REDACTED]-RASL<sup>167</sup>SLEH<sub>6</sub>...RRLL<sup>186</sup>RALEH<sub>6</sub>...RQVA<sup>194</sup>QTLEH<sub>6</sub>

FIG. 24A

PS1 PS2 PS3 PS1 PS2 PS3

kDa

21 →

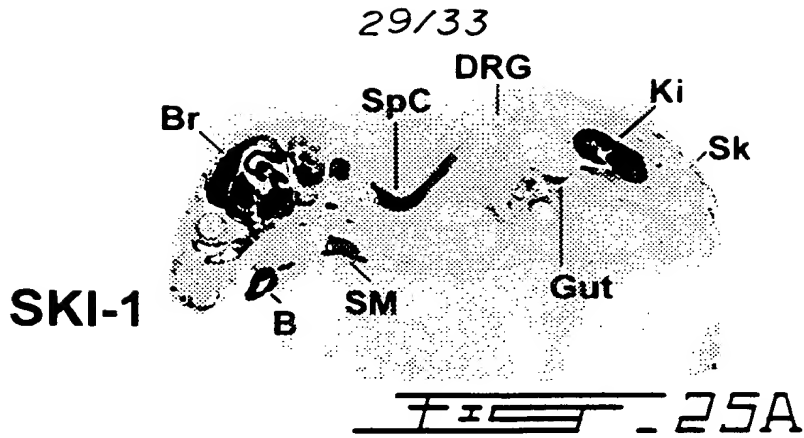


Ab P

Western

Coomassie

FIG. 24B



SKI-1

APP

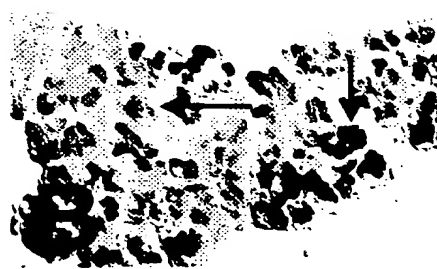
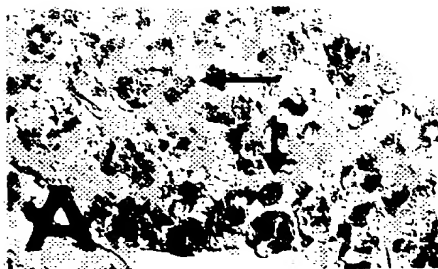


FIG. 26A

FIG. 26B

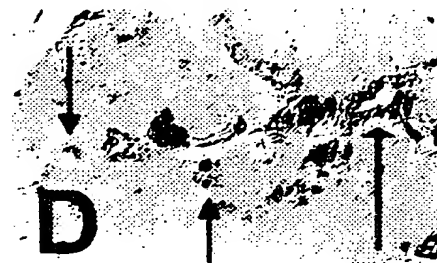
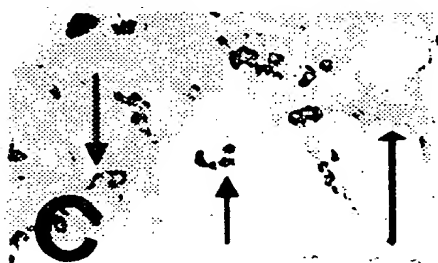
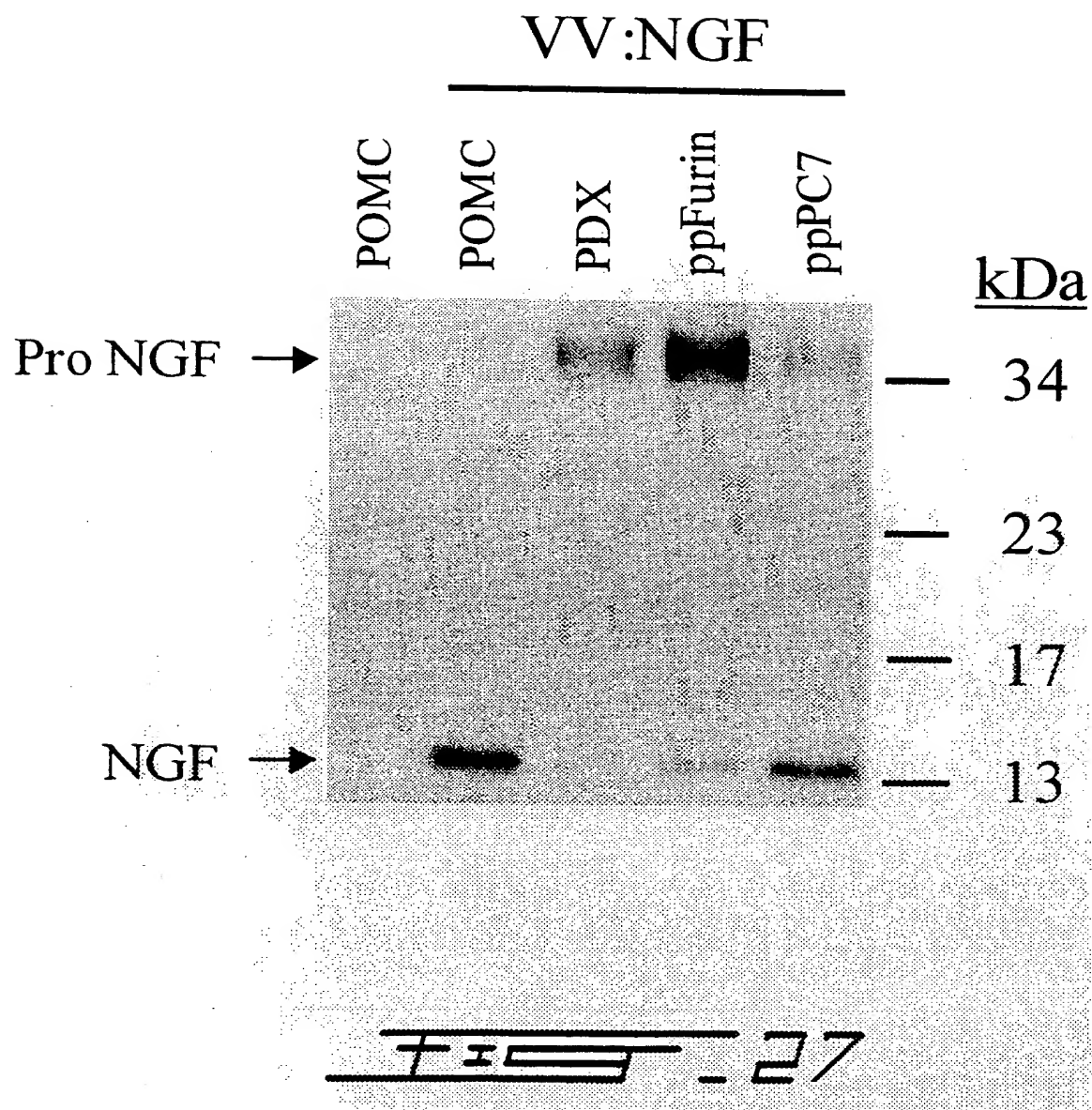


FIG. 26C

FIG. 26D

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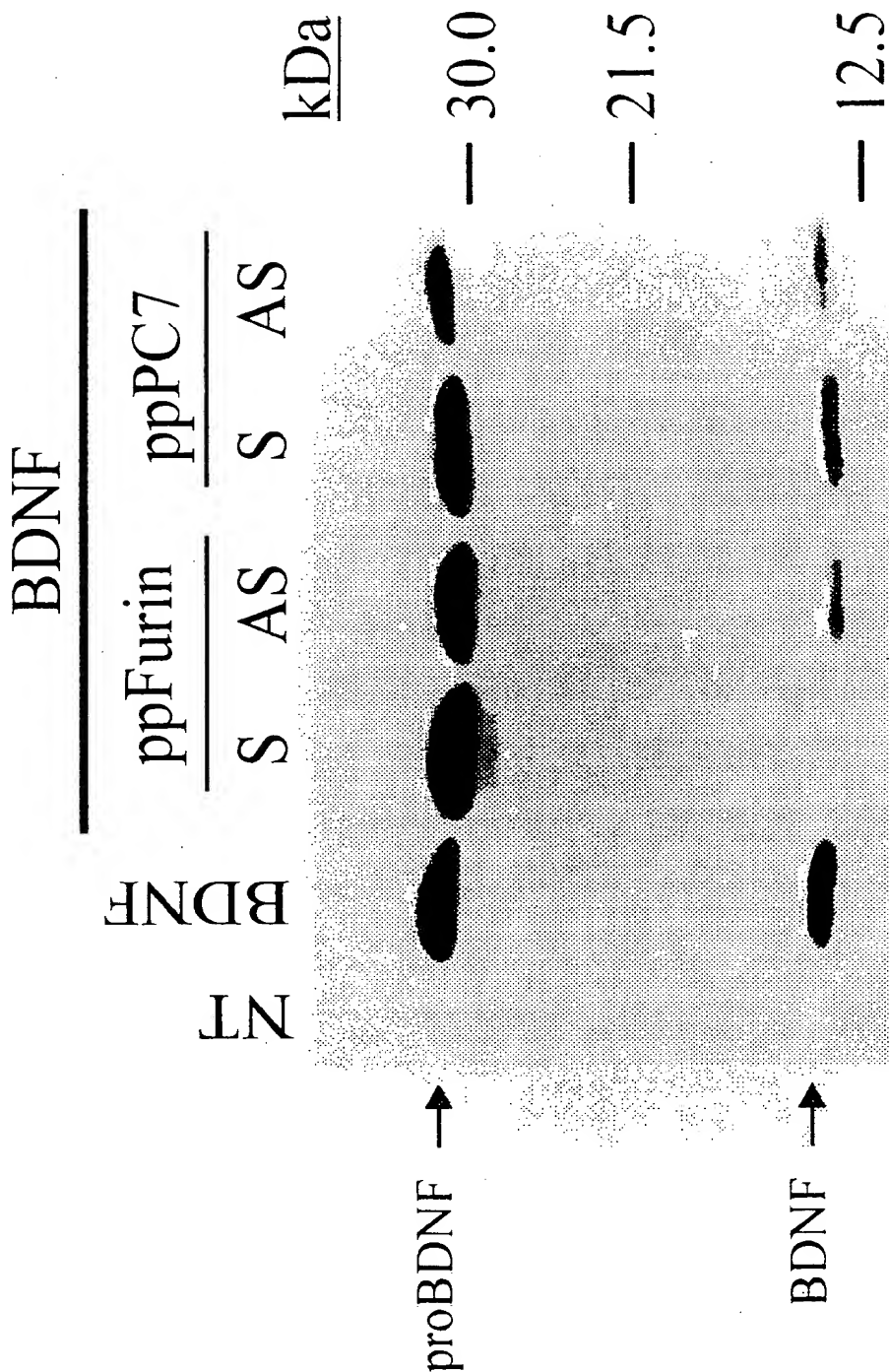
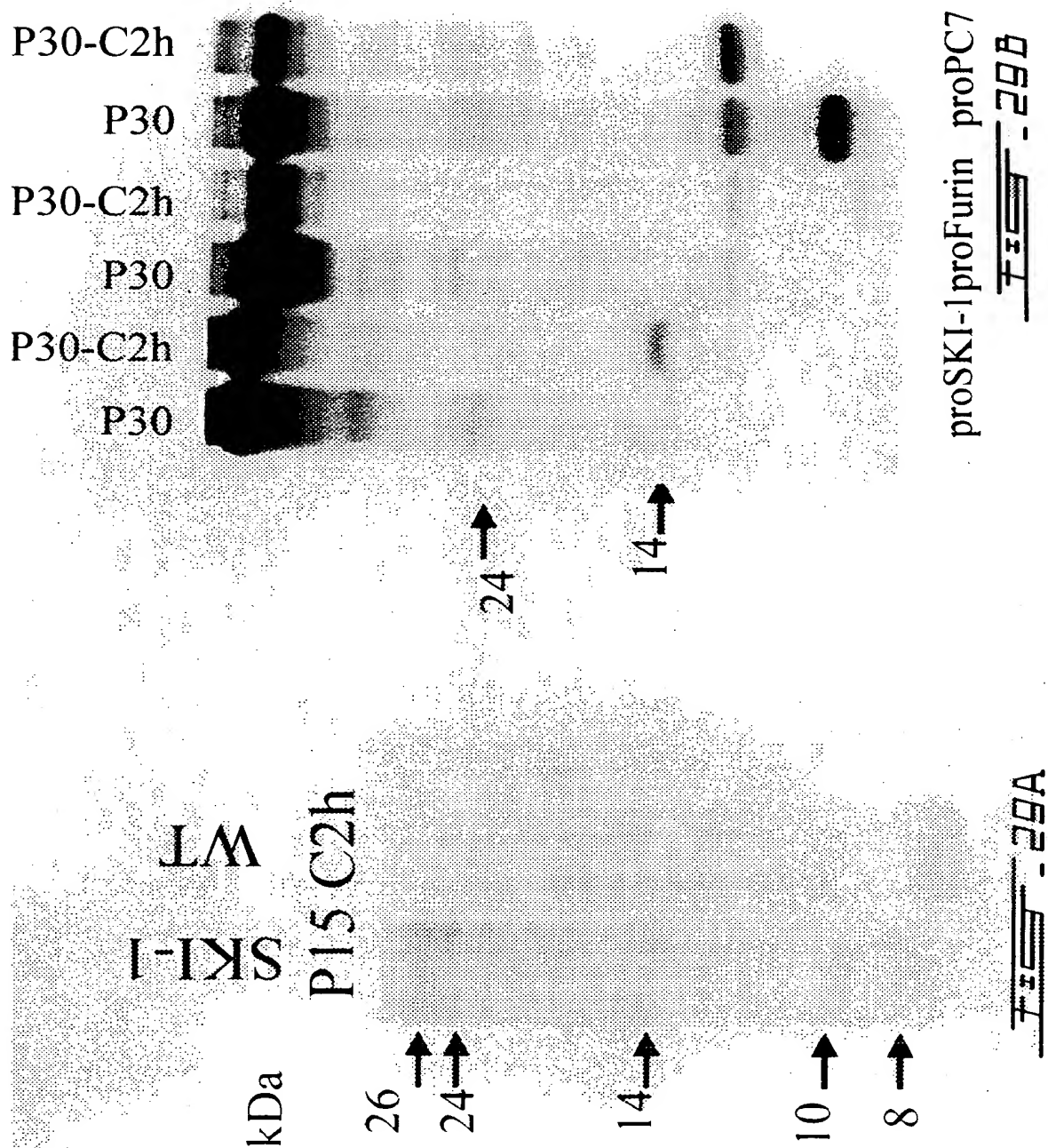


FIG. 28

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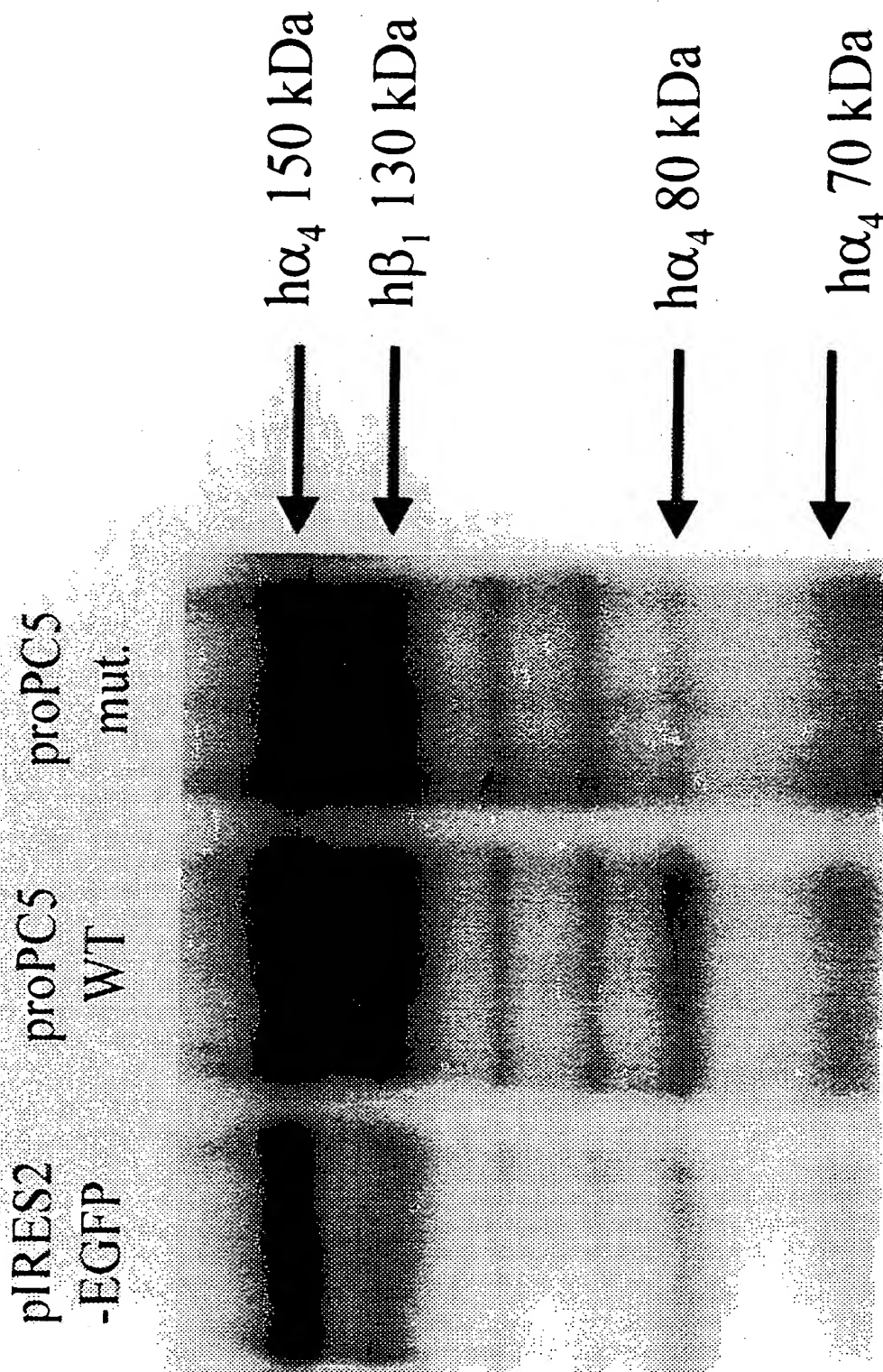


FIG. 30